RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:	10/568.73 <u>7A</u>
Source:	1FWO.
Date Processed by STIC:	. 2/1/07

ENTERED



IFWO

RAW SEQUENCE LISTING DATE: 02/01/2007
PATENT APPLICATION: US/10/568,737A TIME: 08:37:20

Input Set : F:\436uspc.app.txt

```
4 <110> APPLICANT: Rioux, Stephane
             Martin, Denis
     5
     6
             Hamel, Josee
             Brodeur, Bernard R.
     7
     8
             Rheault, Patrick
     9
             Droun, Nathalie
    11 <120> TITLE OF INVENTION: POLYPEPTIDES OF STREPTOCOCCUS PYOGENES
    13 <130> FILE REFERENCE: 484112.436USPC
    15 <140> CURRENT APPLICATION NUMBER: 10/568,737A
C--> 16 <141> CURRENT FILING DATE: 2006-02-15
    18 <150> PRIOR APPLICATION NUMBER: PCT/CA2004/001510
     19 <151> PRIOR FILING DATE: 2004-08-16
    21 <150> PRIOR APPLICATION NUMBER: US 60/495,094
    22 <151> PRIOR FILING DATE: 2003-08-15
    24 <160> NUMBER OF SEQ ID NOS: 46
     26 <170> SOFTWARE: PatentIn version 3.3
    28 <210> SEO ID NO: 1
     29 <211> LENGTH: 537
     30 <212> TYPE: DNA
    31 <213> ORGANISM: Streptococcus pyogenes
    33 <400> SEQUENCE: 1
                                                                               60
     34 atgtcacgta ttggtaataa agtaattact atgcctgcag gcgttgaatt aacaaataac
     35 aacaatgtta ttactgttaa aggccctaaa ggcgaactca ctcgtgagtt caacaaaaat
                                                                              120
     36 attgaaatca aagttgaagg gactgaaatc acagttgtac gtcctaacga ctcaaaagaa
                                                                              180
     37 atgaaaacaa tccatggtac aacccgtgct aacttgaata acatggttgt aggtgtttct
                                                                              240
     38 gaaggtttca aaaaagatct tgaaatgaag ggtgtcggtt accgtgctca acttcaaggt
                                                                              300
     39 actaaacttg tcctttcagt aggtaaatct caccaagacg aagttgaagc tccagaagga
                                                                              360
     40 attactttca ctgttgctaa cccaacttca atctcagttg aaggaatcaa caaagaagtt
                                                                              420
                                                                              480
     41 gttggtcaaa cagctgctta catccgtagc ttgcgttcac cagagcctta caaaggcaaa
     42 gggatccgtt acgttggtga atacgtacgc cttaaagaag gtaaaacagg taaataa
                                                                              537
     45 <210> SEQ ID NO: 2
     46 <211> LENGTH: 178
     47 <212> TYPE: PRT
     48 <213> ORGANISM: Streptococcus pyogenes
     50 <400> SEQUENCE: 2
     51 Met Ser Arg Ile Gly Asn Lys Val Ile Thr Met Pro Ala Gly Val Glu
                                            10
                        5
     53 Leu Thr Asn Asn Asn Val Ile Thr Val Lys Gly Pro Lys Gly Glu
                                        25
     54
                    20
     55 Leu Thr Arg Glu Phe Asn Lys Asn Ile Glu Ile Lys Val Glu Gly Thr
                                    40
                35
     57 Glu Ile Thr Val Val Arg Pro Asn Asp Ser Lys Glu Met Lys Thr Ile
```

RAW SEQUENCE LISTING DATE: 02/01/2007
PATENT APPLICATION: US/10/568,737A TIME: 08:37:20

Input Set : F:\436uspc.app.txt

```
59 His Gly Thr Thr Arg Ala Asn Leu Asn Asn Met Val Val Gly Val Ser
60 65
61 Glu Gly Phe Lys Lys Asp Leu Glu Met Lys Gly Val Gly Tyr Arg Ala
                                        90
                   85
63 Gln Leu Gln Gly Thr Lys Leu Val Leu Ser Val Gly Lys Ser His Gln
               100
64
65 Asp Glu Val Glu Ala Pro Glu Gly Ile Thr Phe Thr Val Ala Asn Pro
                                                    125
                               120
66
67 Thr Ser Ile Ser Val Glu Gly Ile Asn Lys Glu Val Val Gly Gln Thr
       130
                           135
68
69 Ala Ala Tyr Ile Arg Ser Leu Arg Ser Pro Glu Pro Tyr Lys Gly Lys
                       150
                                            155
71 Gly Ile Arg Tyr Val Gly Glu Tyr Val Arg Leu Lys Glu Gly Lys Thr
                                        170
72
                   165
73 Gly Lys
76 <210> SEQ ID NO: 3
77 <211> LENGTH: 1269
78 <212> TYPE: DNA
79 <213> ORGANISM: Streptococcus pyogenes
81 <400> SEQUENCE: 3
82 atgtttcagt taagaaaaaa aatgacgcgc aaacaattag ccttgttgag tgctggagtg
                                                                          60
83 ttgacctgtg tggttggtgg tagctacttg ataatgaacc atcaacaaca agaaattgtc
                                                                          120
                                                                         180
84 tctagtgtca acaaagtaaa agccttaacc ataaaagaag ccatggaaca aggaaaagat
                                                                         240
85 atcagettga cettagetgg egaagtaaca getaacaaca geageaaagt caaaategae
                                                                          300
86 tcaagtaaag gagaagtcaa agaggtcttt gttaaaaaag gcgatgttgt caaagtagga
                                                                          360
87 caaccettgt ttagetatga aacgteacag eggttaaegg eteaaagtte agaatttgat
                                                                          420
88 gttcaaacca aagccaatca gctccaagtt gctaaaacca atgcagcatt gaagtgggaa
89 acctacaatc gcaaggtcaa tgaaatcaac accctaaaat ctcgctacaa cactgcacca
                                                                          480
                                                                          540
90 qatqaqaqct tactaqaqca aattcgcagc gcagaagaca gtgtatccca agcactaagc
                                                                          600
91 gatgccaaaa cagcagatag cgatgtcaaa accgctcaaa tcgaactcga taaagctaat
92 gctactgcca caacggaaaa aggtaaacta gagtatgaca ccgttaagtc agacaccgca
                                                                          660
                                                                          720
93 ggaaccattg ttagtctaaa tactgatttg ccaaatcaat caaaatccaa aaaagaaaat
                                                                          780
94 gaaactttta tggaaattat cgacaaatca aaaatgttag tcaaaggtaa cattagtgaa
                                                                          840
95 tttgaccgtg acaagttaaa aatcggtcaa aaagtcgaag tgattgaccg caaagacaac
96 tctaaaaaat ggactggaaa agtaacccaa gttggcaacc tcaaagcaga ggaaaaaggc
                                                                          900
                                                                          960
97 caaggtcaag gccaaggtgg caatgaccaa caagataatc caaaccaagc aaaattccct
98 tatgttattg aacttgacca atcagacaag cagccactca ttggctcaca cacctatgtt
                                                                         1020
                                                                         1080
99 aatgtgctca acaatgttcc agaagctggc aagatcgtat tgaaagaaac ctttacaatg
100 gcagaaaatg gaaaaaccta tgtgtggaaa gttgataaaa acaaggtcaa aaaacaagaa
                                                                          1140
                                                                          1200
101 atcaagacta agcccttctc aaaaggttat gttgaggtaa caagtggctt gactatgcaa
                                                                          1260
102 gataagattg ctcagccgct tcctggcatg aaagacggta tggaggtagg aagtattgtt
                                                                          1269
103 aaaccttaa
106 <210> SEQ ID NO: 4
107 <211> LENGTH: 422
108 <212> TYPE: PRT
109 <213> ORGANISM: Streptococcus pyogenes
111 <400> SEQUENCE: 4
112 Met Phe Gln Leu Arg Lys Lys Met Thr Arg Lys Gln Leu Ala Leu Leu
113 1
```

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/568,737A

DATE: 02/01/2007
TIME: 08:37:20

Input Set : F:\436uspc.app.txt

114 115	Ser	Ala	Gly	Val 20	Leu	Thr	Cys	Val	Val 25	Gly	Gly	Ser	Tyr	Leu 30	Ile	Met
116	Asn	His	Gln	Gln	Gln	Glu	Ile	Val		Ser	Val	Asn	Lys	Val	Lys	Ala
117			35					40					45			
118	Leu	Thr	Ile	Lys	Glu	Ala	Met	Glu	Gln	Gly	Lys	Asp	Ile	Ser	Leu	Thr
119		50					55					60				
120	Leu	Ala	Gly	Glu	Val	Thr	Ala	Asn	Asn	Ser	Ser	Lys	Val	Lys	Ile	Asp
121	65					70					75					80
122	Ser	Ser	Lys	Gly	Glu	Val	Lys	Glu	Val	Phe	Val	Lys	Lys	Gly	Asp	Val
123					85					90					95	
124	Val	Lys	Val	Gly	Gln	Pro	Leu	Phe	Ser	Tyr	Glu	Thr	Ser	Gln	Arg	Leu
125				100					105					110		
126	Thr	Ala	Gln	Ser	Ser	Glu	Phe	Asp	Val	Gln	Thr	Lys	Ala	Asn	Gln	Leu
127			115					120					125			
128	Gln	Val	Ala	Lys	Thr	Asn	Ala	Ala	Leu	Lys	Trp	Glu	Thr	Tyr	Asn	Arg
129		130					135					140				
130	Lys	Val	Asn	Glu	Ile	Asn	Thr	Leu	Lys	Ser	Arg	Tyr	Asn	Thr	Ala	Pro
131	145					150					155					160
132	Asp	Glu	Ser	Leu	Leu	Glu	Gln	Ile	Arg	Ser	Ala	Glu	Asp	Ser	Val	Ser
133					165				•	170					175	
134	Gln	Ala	Leu	Ser	Asp	Ala	Lys	Thr	Ala	Asp	Ser	Asp	Val	Lys	Thr	Ala
135				180					185					190		
136	Gln	Ile	Glu	Leu	Asp	Lys	Ala	Asn	Ala	Thr	Ala	Thr		Glu	Lys	Gly
137			195					200					205			_
138	Lys	Leu	Glu	Tyr	Asp	Thr	Val	Lys	Ser	Asp	Thr	Ala	Gly	Thr	Ile	Val
139		210					215					220				_
140	Ser	Leu	Asn	Thr	Asp	Leu	Pro	Asn	Gln	Ser	Lys	Ser	Lys	Lys	Glu	Asn
141	225					230					235					240
142	Glu	Thr	Phe	Met	Glu	Ile	Ile	Asp	Lys	Ser	Lys	Met	Leu	Val	Lys	GLY
143					245					250		_			255	
144	Asn	Ile	Ser	Glu	Phe	Asp	Arg	Asp		Leu	Lys	Ile	Gly		Lys	Val
145				260					265					270	_	** - 7
146	Glu	Val	Ile	Asp	Arg	Lys	Asp		Ser	Lys	Lys	Trp	Thr	GLY	ьys	Val
147			275					280			_		285	~1	41	G 3
148	Thr		Val	Gly	Asn	Leu		Ala	Glu	GIu	Lys		GIn	GIY	GIII	Gly
149		290					295	_	_		_	300		T	Dha	Dro
150	Gln	Gly	Gly	Asn	Asp		Gln	Asp	Asn	Pro		GIn	Ата	ьys	Pile	Pro 320
151	305	_	_	_	_	310		_	_		315	D	T	71 0	C1	
	Tyr	Val	Ile	Glu	Leu	Asp	GIn	Ser	Asp	ьys	GIN	Pro	ьeu	116	GTA	Ser
153					325		_	_		330		a 1	7 J _	C1	335	
		Thr	Tyr			Val	Leu	Asn			Pro	GIU	Ald	GTÀ	цуб	Ile
155				340					345		•	a1	T	350		7727
		Leu			Thr	Phe	Thr			GIU	Asn	GIY	гуs	1111	тўт	Val
157			355		_	_	_	360		.	~1	~ 1	365		Thr.	Tarc
				Asp	ьуs	Asn			ьys	ьys	GIN	GIU	тте	пÀг	1117	Lys
159	_	370		_	~-	_	375		77-7	mh		380		ሞኮ∽	Mot	Gln
			Ser	Lys	Gly		vai	GIU	val	ınr		GTĀ	ьeu	. 1111	MEC	Gln 400
161	385	_			0 3	390	T	D	~1	. Ma-	395	7 ~~	روايي	. M≏+	Glu	
162	Asp	Lys	He	Ala	GIn	Pro	ьeu	Pro	GIĀ	Mec	ьys	ASP	дту	MEL	GIU	Val

RAW SEQUENCE LISTING DATE: 02/01/2007
PATENT APPLICATION: US/10/568,737A TIME: 08:37:20

Input Set : F:\436uspc.app.txt

```
410
                                                             415
163
164 Gly Ser Ile Val Lys Pro
                420
167 <210> SEQ ID NO: 5
168 <211> LENGTH: 885
169 <212> TYPE: DNA
170 <213> ORGANISM: Streptococcus pyogenes
172 <400> SEQUENCE: 5
173 atgataaaac gatgtaaagg aattggtcta gccttaatgg ccttcttttt ggtagcttgt
                                                                           60
174 gtgaatcagc accctaaaac ggctaaagag actgaacagc agagaattgt agccacttcg
                                                                          120
175 gttgctgtgg ttgatatctg tgaccgttta aatttagacc tcgttggggt ttgtgatagt
                                                                          180
176 aaattatata cccttcctaa acgctatgat gctgttaagc gtgtgggttt acccatgaat
                                                                          240
177 cctgatatag agttgattgc ttctttgaaa ccaacttgga ttttgagtcc caattcttta
                                                                          300
178 caagaagatt tggaacccaa gtatcaaaaa ttggatactg agtatggttt tttgaactta
                                                                          360
179 cgaagtgttg agggcatgta ccagtccatt gatgatttag ggaacctttt ccaacgtcaa
                                                                          420
180 caagaagcaa aagaattgcg ccagcaatac caggactatt atcgtgcttt ccaagctaaa
                                                                          480
181 cgtaagggga agaaaaagcc taaagtgctt attcttatgg gcttgccagg tagttatttg
                                                                          540
182 gtggcgacga accaatctta tgtagggaat cttttggact tggcaggtgg tgagaatgtt
                                                                          600
                                                                          660
183 tatcagtcag atgagaaaga atttctatca gctaatcctg aagacatgct ggctaaggag
184 cctgacttga ttttacgaac agctcatgcc attccagaca aggtaaaagt gatgtttgac
                                                                          720
                                                                          780
185 aaagaatttg ctgaaaatga tatttggaaa cattttacgg cagtcaagga agggaaagtc
186 tatgatttgg acaataccct gtttggcatg agtgctaaat tgaactaccc agaagccttg
                                                                          840
                                                                           885
187 gacaccttaa cacagctttt tgaccacgtg ggagatcatc cgtaa
189 <210> SEQ ID NO: 6
190 <211> LENGTH: 294
191 <212> TYPE: PRT
192 <213> ORGANISM: Streptococcus pyogenes
194 <400> SEQUENCE: 6
195 Met Ile Lys Arg Cys Lys Gly Ile Gly Leu Ala Leu Met Ala Phe Phe
196 1
197 Leu Val Ala Cys Val Asn Gln His Pro Lys Thr Ala Lys Glu Thr Glu
                                     25
                20
199 Gln Gln Arg Ile Val Ala Thr Ser Val Ala Val Val Asp Ile Cys Asp
                                 40
201 Arg Leu Asn Leu Asp Leu Val Gly Val Cys Asp Ser Lys Leu Tyr Thr
        50
                            55
203 Leu Pro Lys Arg Tyr Asp Ala Val Lys Arg Val Gly Leu Pro Met Asn
                         70
204 65
205 Pro Asp Ile Glu Leu Ile Ala Ser Leu Lys Pro Thr Trp Ile Leu Ser
                    85
                                         90
207 Pro Asn Ser Leu Gln Glu Asp Leu Glu Pro Lys Tyr Gln Lys Leu Asp
                                                         110
                100
                                     105
208
209 Thr Glu Tyr Gly Phe Leu Asn Leu Arg Ser Val Glu Gly Met Tyr Gln
                                                     125
                                 120
210
211 Ser Ile Asp Asp Leu Gly Asn Leu Phe Gln Arg Gln Gln Glu Ala Lys
                            135
213 Glu Leu Arg Gln Gln Tyr Gln Asp Tyr Tyr Arg Ala Phe Gln Ala Lys
                                             155
216 Arg Lys Gly Lys Lys Pro Lys Val Leu Ile Leu Met Gly Leu Pro
```

RAW SEQUENCE LISTING DATE: 02/01/2007 PATENT APPLICATION: US/10/568,737A TIME: 08:37:20

Input Set : F:\436uspc.app.txt

217 165 170 175										
217 165 170 218 Gly Ser Tyr Leu Val Ala Thr Asn Gln Ser Tyr Val Gly Asn Leu Leu										
105										
219 180 185 190 220 Asp Leu Ala Gly Gly Glu Asn Val Tyr Gln Ser Asp Glu Lys Glu Phe										
205										
221 195 200 205 222 Leu Ser Ala Asn Pro Glu Asp Met Leu Ala Lys Glu Pro Asp Leu Ile										
220										
224 Leu Arg Thr Ala His Ala Ile Pro Asp Lys Val Lys Val Met Phe Asp 235 225 230 230 230										
225 225										
226 Lys Glu Phe Ala Glu Asn Asp Ile Trp Lys His Phe Thr Ala Val Lys										
771										
228 Glu Gly Lys Val Tyr Asp Leu Asp Asn Thr Leu Phe Gly Met Ser Ala										
229 260 265 270 The Can Low Pho Agn										
230 Lys Leu Asn Tyr Pro Glu Ala Leu Asp Thr Leu Thr Gln Leu Phe Asp										
231 275 280 285										
232 His Val Gly Asp His Pro										
233 290										
235 <210> SEQ ID NO: 7										
236 <211> LENGTH: 34										
237 <212> TYPE: DNA										
238 <213> ORGANISM: Artificial Sequence										
240 <220> FEATURE:										
241 <223> OTHER INFORMATION: Oligonucleotide primer										
243 <400> SEQUENCE: 7	34									
244 gagaaaatac atatgtcacg tattggtaat aaag										
247 <210> SEQ ID NO: 8										
248 <211> LENGTH: 29										
249 <212> TYPE: DNA .										
250 <213> ORGANISM: Artificial Sequence										
252 <220> FEATURE:	•									
253 <223> OTHER INFORMATION: Oligonucleotide primer										
255 <400> SEQUENCE: 8										
256 ccctcgagtt atttacctgt tttaccttc	29									
259 <210> SEQ ID NO: 9										
260 <211> LENGTH: 31										
261 <212> TYPE: DNA										
262 <213> ORGANISM: Artificial Sequence										
264 <220> FEATURE:										
265 <223> OTHER INFORMATION: Oligonucleotide primer										
267 <400> SEQUENCE: 9	31									
8 aaggatccca tgtcacgtat tggtaataaa g										
1 <210> SEQ ID NO: 10										
272 <211> LENGTH: 38										
23 <212> TYPE: DNA										
74 <213> ORGANISM: Artificial Sequence										
276 <220> FEATURE:										
277 <223> OTHER INFORMATION: Oligonucleotide primer										
279 <400> SEQUENCE: 10										
280 actagtcgac ttatttacct gttttacctt ctttaagg	38									
280 actagragae tractiacet gereracete erreadgy										

VERIFICATION SUMMARY

DATE: 02/01/2007

PATENT APPLICATION: US/10/568,737A

TIME: 08:37:21

Input Set : F:\436uspc.app.txt

Output Set: N:\CRF4\02012007\J568737A.raw

L:16 M:271 C: Current Filing Date differs, Replaced Current Filing Date